



Fig. 7A

Variants:	1	α	β	2	3
RT-PCR product	NO	+	+	NO	+ & -
PCR from LIM1215 lib.	-	+	-	+	NO
RT-PCR product	NO	-	+	NO	+
53.2 cDNA	-	-	-	-	NO

Fig. 7B

sequence "Y" 104-105 bases

GGCCTCCCCGGGGTCCGGCTCCGGCTGGGGTTGAGGGCGGCCGGGGGAACCAG  
GlyLeuProGlyValGlyValArgLeuGlyLeuArgAlaAlaGlyGlyAsnGln  
AlaSerProGlySerAlaSerGlyTrpGly \* GlyArgProGlyGlyThrSer  
ProProArgGlyArgArgProAlaGlyValGluGlyGlyArgGlyGluProAla

CGACATGCGGAGAGCAGCGCAGGCGACTCAGGGCGCTTCCCCCGCAGGTG  
ArgHisAlaGluSerSerAlaGlyAspSerGlyArgPheProArgArg  
AspMetArgArgAlaAlaGlnAlaThrGlnGlyAlaSerProAlaGly  
ThrCysGlyGluGlnArgArgArgLeuArgAlaLeuProProGlnVal

sequence "1" 38 bases

GTGGCTGTGCTTTGGTTAACTTCCTTTTAAACCAGAA  
ValAlaValLeuTrpPheAsnPheLeuPheAsnGlnLys

sequence "2" 36 bases

GTGGATGTGACGGGCGGTACGACACCATCCCCAG  
ValAspValThrGlyAlaTyrAspThrIleProGln

sequence "3" 182 bases

GTCTCTACCTTGACAGACCTCCAGCCGTACATGCGACAGTTCGTGGCTCACCTG  
ValSerThrLeuThrAspLeuGlnProTyrMetArgGlnPheValAlaHisLeu

CAGGAGACCAGCCCGCTGAGGGATGCCGTGTCATCGAGCAGAGCTCCTCCCTG  
GlnGluThrSerProLeuArgAspAlaValValIleGluGlnSerSerSerLeu

AATGAGGCCAGCAGTGGCCTCTTCGACGTCTTCCTACGCTTCATGTGCCACCAC  
AsnGluAlaSerSerGlyLeuPheAspValPheLeuArgPheMetCysHisHis

GCCGTGCGCATCAGGGGCAA  
AlaValArgIleArgGlyLys

partial sequence "2" unknown length

GTGAGCGCACCTGGCCGGAAGTGGAGCCTGTGCCCGGCTGGGGCAGGTGCTGCTGCAG  
Ter

GGCCGTTGCGTCCACCTCTGCTTCCGTGTGGGGCAGGCGACTGCCAATCCCAAAGGGT  
CAGATGCCACAGGGTGCCCTCGTCCCATCTGGGGCTGAGCACAAATGCATCTTTCTG  
TGGGAGTGAGGGTGCCTCACACGGGAGCAGTTTTCTGTGCTATTTTGGTAA...

*Fig. 10A*